



## SEQUENCE LISTING

<110> Soo Young Lee  
Yongwon Choi

<120> Signal Transducer for the TNF Receptor  
Super Family and Uses Thereof

<130> 600-1-198CIP1CON

<140> 09/716,536

<141> 2000-11-20

<150> 60/042,293

<151> 1997-04-01

<150> 60/042,747

<151> 1997-04-07

<150> 08/834,903

<151> 1997-04-07

<160> 16

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 469

<212> PRT

<213> Homo sapiens

<400> 1

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Ser	Arg	Asp	Val	Ala	Ala	Ile	His	Cys	Gly	His	Thr	Phe	His	Leu	Gln
			20					25					30		
Cys	Leu	Ile	Gln	Ser	Phe	Glu	Thr	Ala	Pro	Ser	Arg	Thr	Cys	Pro	Gln
			35				40					45			
Cys	Arg	Ile	Gln	Val	Gly	Lys	Arg	Thr	Ile	Ile	Asn	Lys	Leu	Phe	Phe
			50			55					60				
Asp	Leu	Ala	Gln	Glu	Glu	Asn	Val	Leu	Asp	Arg	Glu	Phe	Leu	Lys	
65				70				75						80	
Asn	Glu	Leu	Asp	Asn	Val	Arg	Ala	Gln	Leu	Ser	Gln	Lys	Asp	Lys	Glu
				85				90					95		
Lys	Arg	Asp	Ser	Gln	Val	Ile	Ile	Asp	Thr	Leu	Arg	Asp	Thr	Leu	Glu
			100					105					110		
Glu	Arg	Asn	Ala	Thr	Val	Val	Ser	Leu	Gln	Gln	Ala	Leu	Gly	Lys	Ala
			115				120					125			
Glu	Met	Leu	Cys	Ser	Thr	Leu	Lys	Lys	Gln	Met	Lys	Tyr	Leu	Glu	Gln
						135					140				
Gln	Gln	Asp	Glu	Thr	Lys	Gln	Ala	Gln	Glu	Glu	Ala	Gly	Arg	Leu	Arg
145					150				155					160	
Ser	Lys	Met	Lys	Thr	Met	Glu	Gln	Ile	Glu	Leu	Leu	Leu	Gln	Ser	Gln
				165					170					175	
Leu	Pro	Glu	Val	Glu	Glu	Met	Ile	Arg	Asp	Met	Gly	Val	Gly	Gln	Ser

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TECH CENTER 1600/2900





1				5					10					15		
Asn	Val	Leu	Asp	Arg	Glu	Phe	Leu	Lys	Asn	Glu	Leu	Asp	Asn	Val	Arg	
			20					25					30			
Ala	Gln	Leu	Ser	Gln	Lys	Asp	Lys	Glu	Lys	Arg	Asp	Ser	Gln	Val	Ile	
		35					40					45				
Ile	Asp	Thr	Leu	Arg	Asp	Thr	Leu	Glu	Glu	Arg	Asn	Ala	Thr	Val	Val	
	50					55					60					
Ser	Leu	Gln	Gln	Ala	Leu	Gly	Lys	Ala	Glu	Met	Leu	Cys	Ser	Thr	Leu	
65					70				75						80	
Lys	Lys	Gln	Met	Lys	Tyr	Leu	Glu	Gln	Gln	Gln	Asp	Glu	Thr	Lys	Gln	
				85				90						95		
Ala	Gln	Glu	Glu	Ala	Gly	Arg	Leu	Arg	Ser	Lys	Met	Lys	Thr	Met	Glu	
			100					105					110			
Gln	Ile	Glu	Leu	Leu	Leu	Gln	Ser	Gln	Leu	Pro	Glu	Val	Glu	Glu	Met	
		115					120					125				
Ile	Arg	Asp	Met	Gly	Val	Gly	Gln	Ser	Ala	Val	Glu	Gln	Leu	Ala	Val	
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Tyr	Cys	Val	Ser	Leu	Lys	Lys	Glu	Tyr	Glu	Asn	Leu	Lys	Glu	Ala	Arg	
145					150				155						160	
Lys	Ala	Ser	Gly	Glu	Val	Ala	Asp	Lys	Leu	Arg	Lys	Asp	Leu	Phe	Ser	
			165					170					175			
Ser	Arg	Ser	Lys	Leu	Gln	Thr	Val	Tyr	Ser	Glu	Leu	Asp	Gln	Ala	Lys	
			180					185					190			
Leu	Glu	Leu	Lys	Ser	Ala	Gln	Lys	Asp	Leu	Gln	Ser	Ala	Asp	Lys	Glu	
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	210					215					220					

<210> 4

<211> 220

<212> PRT

<213> Mus musculus

<400> 4

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			20					25					30			
Ala	Gln	Leu	Ser	Gln	Lys	Asp	Arg	Glu	Lys	Arg	Asp	Ser	Gln	Ala	Ile	
		35					40					45				
Ile	Asp	Thr	Leu	Arg	Asp	Thr	Leu	Glu	Glu	Arg	Asn	Ala	Thr	Val	Glu	
	50					55					60					
Ser	Leu	Gln	Asn	Ala	Leu	Asn	Lys	Ala	Glu	Met	Leu	Cys	Ser	Thr	Leu	
65					70				75						80	
Lys	Lys	Gln	Met	Lys	Phe	Leu	Glu	Gln	Arg	Gln	Asp	Glu	Thr	Lys	Gln	
				85				90						95		
Ala	Arg	Glu	Glu	Ala	His	Arg	Leu	Lys	Cys	Lys	Met	Lys	Thr	Met	Glu	
			100					105					110			
Gln	Ile	Glu	Leu	Leu	Leu	Gln	Ser	Gln	Arg	Ser	Glu	Val	Glu	Glu	Met	
		115					120					125				
Ile	Arg	Asp	Met	Gly	Val	Gly	Gln	Ser	Ala	Val	Glu	Gln	Leu	Ala	Val	
	130					135					140					
Tyr	Cys	Val	Ser	Leu	Lys	Lys	Glu	Tyr	Glu	Asn	Leu	Lys	Glu	Ala	Arg	
145					150				155						160	
Lys	Ala	Thr	Gly	Glu	Leu	Ala	Asp	Arg	Leu	Lys	Lys	Asp	Leu	Val	Ser	
				165				170					175			

Ser Arg Ser Lys Leu Lys Thr Leu Asn Thr Glu Leu Asp Gln Ala Lys  
180 185 190  
Leu Glu Leu Arg Ser Ala Gln Lys Asp Leu Gln Ser Ala Asp Gln Glu  
195 200 205  
Ile Thr Ser Leu Arg Lys Lys Ser Asp Asp Pro Pro  
210 215 220

<210> 5  
<211> 51  
<212> PRT  
<213> Homo sapiens

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Val Ala Ala Met Asp Cys Gly His Thr Phe His Leu Gln Cys Leu Ile  
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Gln Ser Phe Glu Thr Ala Pro Ser Arg Thr Cys Pro Gln Cys Arg Ile  
35 40 45  
Gln Val Gly  
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<210> 6  
<211> 51  
<212> PRT  
<213> Mus musculus

<400> 6  
Leu Ser Leu Cys Thr Ile Cys Ser Asp Phe Phe Asp His Ser Arg Asp  
1 5 10 15  
Val Ala Ala Ile His Cys Gly His Thr Phe His Leu Gln Cys Leu Ile  
20 25 30  
Gln Trp Phe Glu Thr Ala Pro Ser Arg Thr Cys Pro Gln Cys Arg Ile  
35 40 45  
Gln Val Gly  
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<210> 7  
<211> 2007  
<212> DNA  
<213> Homo sapiens

<400> 7  
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tgcactatct gctccgactt cttcgatcac tcccgcgacg tggccgccat ccactgcggc 180  
cacaccttcc acttgcagtg cctaattcag tcctttgaga cagcaccaag tcggacctgc 240  
ccacagtgcc gaatccaggt tggcaaaaga accattatca ataagctctt ctttgatctt 300  
gcccaggagg aggagaatgt cttggatcga gaattcttaa agaatgaact ggacaatgtc 360  
agagcccagc tttcccagaa agacaaggag aaacgagaca gccaggatcat catcgacact 420  
ctgcgggata cgctggaaga acgcaatgct actgtggtat ctctgcagca ggccttgggc 480  
aaggccgaga tgctgtgctc cacactgaaa aagcagatga agtacttaga gcagcagcag 540  
gatgagacca aacaagcaca agaggaggcg ggccgggtca ggagcaagat gaagaccatg 600  
gagcagattg agcttctact ccagagccag ctccctgagg tggaggagat gatccgagac 660

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gagtacgaga	atctaaaaga	ggcacggaag	gcctcagggg	aggtggctga	caagctgagg	780
aaggatttgt	tttcctccag	aagcaagttg	cagacagtct	actctgaatt	ggatcaggcc	840
aagttagaac	tgaagtcagc	ccagaaggac	ttacagagtg	ctgacaagga	aatcatgagc	900
ctgaaaaaga	agctaacgat	gctgcaggaa	accttgaacc	tgccaccagt	ggccagttag	960
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cggccatcct	tccgtgatga	tattgatctc	aatgctacct	ttgatgtgga	tactccccca	1080
gcccggccct	ccagctccca	gcatggttac	tacgaaaaac	tttgcctaga	gaagtcacac	1140
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tcactgggtg	gccagagctg	tgcaggagag	ccagatgagg	aactggttgg	tgccttcctt	1260
atTTTTgtcc	ggaatgccat	cctaggccag	aaacagccca	aaaggcccag	gtcagagtcc	1320
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<210> 8

<211> 1975

<212> DNA

<213> Mus musculus

<400> 8

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ctctctctgt	gcactatctg	ctccgacttc	ttcgatcact	cccgtgacgt	ggctgccatc	180
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gacagcgtca	aagctcagct	ttcccagaaa	gacagggaga	aacgggacag	ccaggccatt	420
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cagcggcagg	atgagaccaa	acaagctcgg	gaggaggccc	accgactcaa	gtgcaagatg	600
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ctcaagaaag	agtatgagaa	tctgaaggaa	gctcggaagg	ccacagggga	actggctgac	780
aggttgaaga	aggatttggg	gtcctctagg	agcaagttga	agactctcaa	cactgagctg	840
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cgaacaaaat	tcattccagcc	tagggacaca	accattatcc	gaccagtgcc	tgtaagtcc	1440
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ctggatacct	tcttatgtca	gtgaacgggtg	accagagtga	tgtttgcaat	tagtgggcca	1560
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gtggatggga gtgctggagg atcctatgca ggctggagga ccctgcgctt gaactcctgc 1860
ctgcctccag cttattgctt gaaattatgg ggtgaggtgg tgatagggaa aggttgggga 1920
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<210> 9
<211> 47
<212> PRT
<213> Artificial Sequence

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<220>
<223> fragment

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Ser Ser Gly Pro Gln Asn Cys Ala Ala Cys Val Tyr Glu Gly Leu
      35             40             45

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<210> 10
<211> 46
<212> PRT
<213> Artificial Sequence

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<220>
<223> fragment

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Lys Tyr Lys Cys Glu Lys Cys Arg Leu Val Leu Cys Asn Pro Lys Gln
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Thr Glu Cys Gly His Arg Phe Cys Glu Ser Cys Met Ala Ala Leu Leu
             20             25             30
Ser Ser Ser Ser Pro Lys Cys Thr Ala Cys Gln Glu Ser Ile
      35             40             45

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<210> 11
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<212> PRT
<213> Artificial Sequence

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<220>
<223> fragment

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             20             25             30
Arg Lys Cys Pro Ile Cys Gly Arg Gly Thr Ile
      35             40

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<210> 12  
<211> 47  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> fragment

<400> 12  
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Glu Pro Cys Gly His Leu Met Cys Thr Ser Cys Leu Thr Ser Trp Gln  
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Glu Ser Glu Gly Gln Gly Cys Pro Phe Cys Arg Cys Glu Ile Lys  
35 40 45

<210> 13  
<211> 48  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> fragment

<400> 13  
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1 5 10 15  
Thr Lys Glu Cys Leu His Arg Phe Cys Ser Asp Cys Ile Val Thr Ala  
20 25 30  
Leu Arg Ser Gly Asn Lys Glu Cys Pro Thr Cys Arg Lys Lys Leu Val  
35 40 45

<210> 14  
<211> 47  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> fragment

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1 5 10 15  
Ile Glu Cys Gly His Ser Phe Cys Gln Glu Cys Ile Ser Gln Val Gly  
20 25 30  
Lys Gly Gly Gly Ser Val Cys Ala Val Cys Arg Gln Arg Phe Leu  
35 40 45

<210> 15  
<211> 50  
<212> PRT  
<213> Artificial Sequence



